Two new tetraploid quillworts species, Iso ëtes longpingii and

I. xiangfei from China (Iso ëtaceae)

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Abstract: Two new tetraploids quillworts species, *Iso äes longpingii* and *I. xiangfei*, from China are described and illustrated. *I. longpingii* is a fully submerged plant found in a small pond of Hunan Province. It is morphologically similar to *I. sinensis*, but differs in its small and tuberculate-cristate megaspores and soft slender leaves that grow up to 60 cm. It is also similar to hexaploid *I. orientalis*, but differs in its 44 chromosomes and tuberculate-cristate megaspores. *I. xiangfei* is most similar to diploid *I. yunguiensis* in megaspore ornamentation, but differs in its microspore ornamentation, oblong sporangium, and 44 chromosomes. There are a few individuals of *I. longpingii* found in Ningxiang County of Hunan Province, and *I. xiangfei* is distributed in the wetlands of Tongdao and Huitong counties of Hunan Province. Because of their limited geographic range, small populations, fewer individuals and disturbed habitats, *I. longpingii* and *I. xiangfei* are ecaluated as critically endangered (CR) and vulnerable (VU) category, respectively, according to the IUCN Red List criteria. A key to all the current known Chinese quillworts is also provided for further taxonomic identification and conservation of these rare and endangered plants in China.

Keywords: new taxon, *Iso äes*, Lycopods, polyploid, heterosporous, aquatic plants

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中国水韭属两个四倍体新种

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摘要:水韭属(Iso äes)是起源最为古老的水生维管植物,全属物种均被列为国家一级重点保护植物。通过对全国水韭属植物的调查和研究,发现不同产地的四倍体植株在形态上存在显著差异。基于形态学、孢粉学和细胞学证据,将分布于中国湖南省长沙地区和怀化地区的四倍体居群分别命名为隆平水韭(Iso äes longpingii)和湘妃水韭(I. xiangfei),并详细描述了其形态特征。隆平水韭形态上与中华水韭相似,但不同之处在于其大孢子具小的瘤状或冠状纹饰,叶细长而柔弱,长达 60 cm;该种也与六倍体东方水韭(I. orientalis)相似,不同之处在于其染色体 44 条,大孢子具瘤状或冠状纹饰。湘妃水韭的大孢子纹饰与二倍体云贵水韭(I. yunguiensis)相似,但在小孢子纹饰、孢子囊形状和染色体数目方面不同。隆平水韭仅少数植株生长于湖南省宁乡市一处池塘,完全沉水生长,而湘妃水韭分布于怀化市通道县和会同县的湿地。由于这两个新种的分布区狭窄,野生居群数量和个体数较少,栖息地环境受到人为干扰,根据 IUCN 红色名录评估标准,将隆平水韭评为极危(CR)等级,湘妃水韭评为易危(VU)等级。编制的中国已知水韭属物种的分种检索表为本属物种的鉴定和保护工作提供了重要参考。

关键词: 新分类群, 水韭属, 石松类植物, 多倍体, 异型孢子, 水生植物

Iso äes L. (1753: 1100) is the only extant genus of the family Iso äaceae, with an extensive fossil record dating from the Devonian Period (Pigg, 2001; Wang et al., 2019) to rapid diversification in the Cenozoic Era (Pereira et al., 2017b, 2021; Wood et al., 2020). The genus contains about 250 species, widely distributed from tropical to subarctic regions and grow in various habitats in most parts of the world (PPGI, 2016; Troia et al., 2016). However, most Iso äes species have restricted habitats and small populations, putting them at risk of extinction (Kang et al., 2005; Liu X et al., 2005; Gentili et al., 2010; Troia et al., 2016; Brunton & Troia, 2018; Singh et al., 2021). Interestingly, hybridization and polyploidization frequently occur in *Iso äes*, and more than half of the known species are polyploids ranging from tetraploid to dodecaploid (Hickey, 1984; Taylor & Hickey, 1992; Liu et al., 2004; Kim et al., 2010; Pereira, 2015; Troia et al., 2016; Grigoryan et al., 2020). Although this ancient genus is widely distributed in various habitats and has a complex evolutionary history, its simple morphology—a tuft of linear sporophylls on the lobed rootstock—is remarkably similar among species, making interspecific identification difficult (Taylor & Hickey, 1992; Troia et al., 2016). Iso äes species are mostly distributed in America, southern Europe, Africa, and Australia, with relatively few species in Asia (Troia et al., 2016; Pereira et al., 2017b, 2021; Brunton & Troia, 2018; Choi et al., 2018; Singh et al., 2021). The Iso äes species diversity in Asia is probably underestimated, and many cryptic species may not be identified (Schafran, 2019).

Characteristics such as habitat, velum, ligule, megaspore size and texture, chromosome count, and DNA sequences are generally employed to distinguish species of *Iso äes* (Hickey, 1986; Pereira, 2015; Troia et al., 2016), however, the characteristics of leaf morphology and anatomy are

influenced by local environmental conditions (Liu et al., 2006; Cavalheiro-Filho et al., 2021; Singh et al., 2021). Several species have recently been described and illustrated around the world (Mora-Olivo et al., 2016; Pereira et al., 2016, 2017a, 2019; Schafran et al., 2016; Li et al., 2019; Lu et al., 2021). In China, seven species—*I. yunguiensis* Q.F. Wang & W.C. Taylor, *I. hypsophila* Hand.-Mazz., *I. taiwanensis* De Vol, *I. shangrilaensis* X. Li & Y.Q. Huang, *I. baodongii* Y. F. Gu, Y. H. Yan & Yi J. Lu, *I. sinensis* T.C. Palmer and *I. orientalis* H. Liu & Q.F. Wang—have been reported (Handel-Mazzetti, 1923; Palmer, 1927; De Vol, 1972; Wang et al., 2002; Liu H et al., 2005; Zhang & Taylor, 2013; Li et al., 2019; Lu et al., 2021).

We conducted several field surveys of the genus in China and collected some tetraploids samples (2n=4x=44) previously considered as *I. sinensis* (2n=4x=44) with morphological and anatomical variation (Liu et al., 2004; Liu et al., 2006; Dai et al., 2020, 2021). By comparing morphological features and spore texture, these samples from Ningxiang and Tongdao counties in Hunan Province were found to be different from *I. sinensis*. In addition, our plastid phylogenomic analysis indicated that these samples formed monophyletic clades, which did not cluster together with *I. sinensis* (Gu et al., unpublished). The sample from Ningxiang County, here named *I. longpingii*, is similar to *I. orientalis* (2n=6x=66) on megaspore ornamentation and that from Tongdao County, here named *I. xiangfei*, is similar to *I. yunguiensis* (2n=2x=22) on megaspore ornamentation. These two new species are described and illustrated in the present study.

1 Materials and Methods

Iso àes species possess prominent spore morphological characteristics, their spore ornamentation terminology established by Hickey (1986) was used in the present study. Megaspores and microspores of samples were examined under a scanning electron microscope. Spores were mounted on a double-sided adhesive tape attached to metal stubs, sputter-coated with platinum, and observed under a field emission scanning electron microscope (Quanta250; FEI, Hillsboro, Oregon, U.S.A.) at 30 kV. Megaspore and microspore size from 3-5 individuals were measured using Photoshop CS5 (Adobe Systems Inc., USA).

To determine the chromosome number, young root tips of the sporophytes were pretreated in a saturated aqueous solution of p-dichlorobenzene for 3–5 h and subsequently fixed in Carnoy's solution (1:3, glacial acetic acid:95% ethanol) for 1 h at 4 °C. Subsequently, the samples were hydrolyzed with a mixture of 3% cellulase and 2.5% pectinase for 10 min at room temperature. They were then stained with carbol fuchsin. The chromosomes of the samples were counted and photographed using a Carl Zeiss Axio Scope A1 photomicroscope (Jena, Germany).

2 Taxonomic Treatment

Iso des longpingii Y.H. Yan, Y.F. Gu & J.P. Shu (Fig. 1, Fig. 3: A-D)

Diagnosis:—*Iso äes longpingii* is similar to *I. sinensis*, but differs in its megaspores small (310–410 μm, mean=350 μm, n=20), sparsely tuberculate-cristate [vs larger (mean 410 μm) densely tuberculate-cristate]and rhizome corms bilobed (vs trilobed in *I. sinensis*). It is also similar to hexaploid *I. orientalis*, but differs in its megaspores tuberculate-cristate, and chromosomes 44 (vs 66 in *I. orientalis*) (Fig. 3, Fig. 4, Table 1).

Type:—CHINA. Hunan Province, Changsha City, Ningxiang County, Yujia'ao Village, 28 °14'27.91" N, 112 °17'44.86" E, alt. 131 m, June 15, 2019. Zhiguo Ou YYH15160 (holotype, PE!; isotype: NOCC!, IBSC!).

Description:—Plants aquatic. Rhizome corms: 2-lobed. Sporophylls: widely spreading, 40–60 in a tuft, white at base, green above, spirally arranged, 20–60 cm long, ca. 1 mm wide, flattened on the adaxial side, rounded on the abaxial side, base flat and alate, peripheral fibrous bundles present, central intrastelar canal 4. Sporangia: basal, obovate, 3.5– 4.5×2.5 –3.0 mm. Ligule: cuniform, 2.3– 2.6×1.1 –1.3 mm. Megaspores: white when dry, earthy yellow when wet, 310–410 μm (mean =350 μm, n=20) in diameter, proximal hemisphere tuberculate, distal hemisphere echinate-cristate. Microspores: gray *en masse*, elliptic, vertical axis length 27–30 μm (mean=29 μm, n=20), surface echinate. Chromosome numbers 2n=4x=44.

Distribution:—Ningxiang County, Changsha City, Hunan Province.

Ecology:—Submerged in a small pond; alt. 131 m.

Etymology:—The specific epithet "longpingii" is derived from the name of the late Prof. Long-Ping Yuan, who devoted his entire life to hybrid rice research in Changsha City (where the species was discovered) and greatly contributed to global food security and poverty alleviation (Wang, 2021).

IUCN Red List category:—A single population of about ten individuals is found in only one location in Yujia'ao Village, Ningxiang County, Hunan Province. *Iso äes longpingii* is critically endangered (CR) [A1a; B2ab(iv)c; C2a(ii); D(2)] according to the Red List Categories and Criteria (Subcommittee, 2019).



A. Habitats; **B.** Microsporangia; **C.** Megasporangia; **D.** Chromosomes (2n=4x=44); **E.** Transection of leaf; **F.** Ligule; **G.** Transection of rhizome (2-lobed).

Fig. 1 Morphological and anatomic characteristics of *Iso äes longpingii* Y.H. Yan,

Y.F. Gu & J.P. Shu

Iso des xiangfei Y.H. Yan, Y.F. Gu & J.P. Shu (Fig. 2, Fig. 4: A-D)

Diagnosis:—*Iso äes xiangfei* is a tetraploid, morphologically similar to the diploid *I. yunguiensis* and the tetraploid *I. sinensis* in megaspore ornamentation, but differs in its less reticulate, lower megaspore ornamentation, and densely low-tuberculate (vs. sparsely tuberculate-echinate) microspores (Fig. 3, Fig. 4, Table 1).

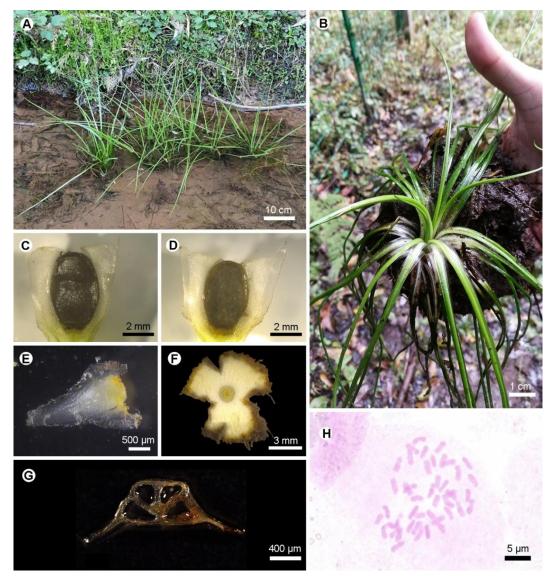
Type:—CHINA. Hunan Province: Huaihua City, Tongdao County, $26^{\circ}14'56.82''$ N, $109^{\circ}50'0.33''$ E, alt. 298 m, December 28, 2019. Juan Yang Fern08928 (holotype: PE!; isotype: NOCC!, IBSC!). **Description:**—Plants aquatic. Rhizome corms 3-lobed. Sporophylls 20–60 in a tuft, widely spreading, white at the base, green above, spirally arranged, 15–35 cm long, 2–3 mm wide, flattened on the adaxial side, rounded on the abaxial side, base flat and alate, peripheral fibrous bundles present, central intrastelar canal 4. Sporangia: basal, oblong, 4.4– 5.0×2.9 –3.3 mm. Ligule: oval triangle, 2.5– 2.7×1.4 –1.6 mm. Megaspores: white when dry, earthy yellow when wet, 390– $450 \mu \text{m}$ (mean= $430 \mu \text{m}$, n=20) in diameter; proximal hemisphere cristate, distal hemisphere reticulate. Microspores: gray *en masse*, elliptic, monolete, vertical axis 26– $28 \mu \text{m}$ (mean= $27 \mu \text{m}$, n=20), surface echinate. Chromosome numbers 2n=4x=44.

Distribution:—Tongdao and Huitong counties, Huaihua City, Hunan Province.

Ecology:—Wetlands and small gullies; alt. 298 m.

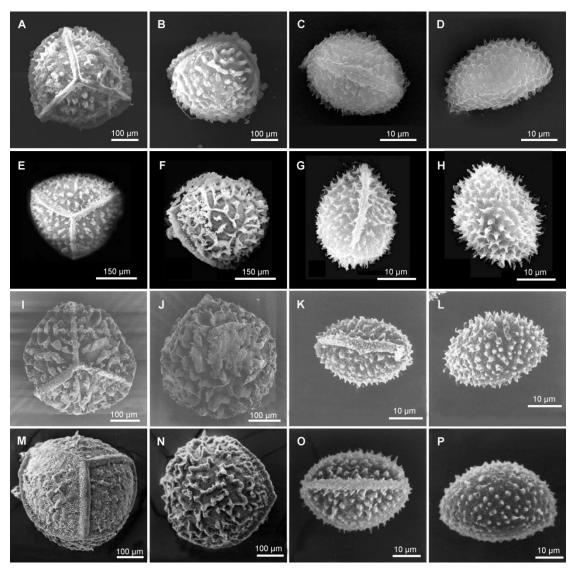
Etymology:—The specific epithet "xiangfei" is derived from the name of Xiangfei, who are two respectable goddesses of Xiangjiang River in Hunan Province, where the aquatic plant was discovered. "Xiang" is the longest River in Hunan Province, and "fei" is the wife of the King in Chinese. Xiangfei, two sisters, married the legendary King Shun at the same time.

IUCN Red List category:—*Iso \(\text{des}\) siangfei* is found in Tongdao and Huitong counties, Hunan Province. There are about 200 individuals, which are well conserved by the local forestry department. It is vulnerable (VU) [B2ab(iii); C2a(i)] according to the Red List Categories and Criteria (Subcommittee, 2019).



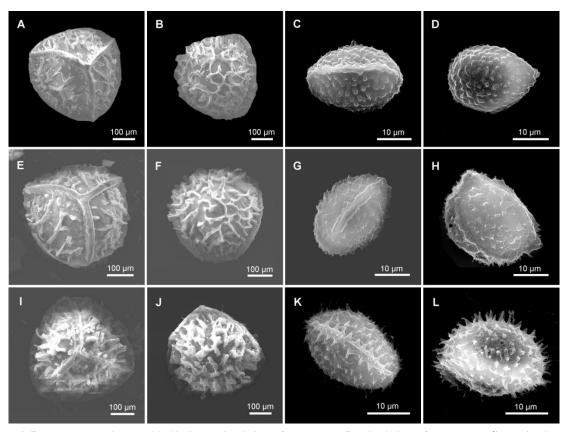
A. Habit; **B**. Plants with wide wing at the base of leaf; **C**. Microsporangia; **D**. Megasporangia; **E**. Ligule; **F**. Transection of rhizome (3-lobed); **G**. Transection of leaf; **H**. Chromosomes.

Fig. 2 Morphological and anatomic characteristics of *Iso &es xiangfei* Y.H. Yan, Y.F. Gu & J.P. Shu



A-D. Iso des longpingii (YYH15160. A. Proximal view of megaspore; B. Distal view of megaspore; C. Proximal view of microspore; D. Distal view of microspore). E-H. I. sinensis from China (cited from Liu et al. 2008. E. Proximal view of megaspore; F. Distal view of megaspore; G. Proximal view of microspore; H. Distal view of microspore). I-L. I. sinensis from Japan (cited from Watanabe et al. 1996. I. Proximal view of megaspore; J. Distal view of megaspore; K. Proximal view of microspore; L. Distal view of microspore). M-P. I. sinensis from Korea (cited from Takamiya 2001. M. Proximal view of megaspore; N. Distal view of megaspore; O. Proximal view of microspore).

Fig. 3 Palynological comparison between Iso ëtes longpingii and I. sinensis



A-D. Iso äes xiangfei (Fern08928. A. Proximal view of megaspore; B. Distal view of megaspore; C. Proximal view of microspore; D. Distal view of microspore). E-H. I. yunguiensis (YYH15163. E. Proximal view of megaspore; F. Distal view of megaspore; G. Proximal view of microspore; H. Distal view of microspore). I-L. I. orientalis (Fern08748. I. Proximal view of megaspore; J. Distal view of megaspore; K. Proximal view of microspore).

Fig. 4 Palynological comparison between Iso äes xiangfei, I. yunguiensis and I. orientalis

Table 1 Spore features of Isoëtes from China

	Chromoso	Megaspore		Microspore	
Species	me number	Ornamentation	Size	Ornamentation	Size
I. hypsophila*	2n=2x=22	Levigate	290–400 μm	Echinate	19–25 μm
			(mean=358 μm)		(mean=22 μm)
I. shangrilaensis	2n=2x=22	Tuberculate-rugulate	207.25–273.09 μm	Echinate-cristate	11.21–23.66 μm
			(mean=244.71 μm)		(mean=19.41 µm)
I. yunguiensis*	2n=2x=22	Cristate-reticulate	340–430 μm	Levigate-granulate	20–25 μm
			(mean=390 μm)		(mean=22 μm)
I. taiwanensis*	2n=2x=22	Tuberculate-cristate	280–340 μm	Echinate	20–28 μm
			(mean=312 μm)		(mean=24 μm)
I. sinensis*	2n=4x=44	Cristate	340–450 μm	Echinate	23–32 μm
			(mean=409 μm)		(mean=28 μm)
I. orientalis*	2n=6x=66	Cristate-reticulate	350–460 μm	Echinate-tuberculate	20–38 μm
			(mean=420 μm)		(mean=34 μm)

I. baodongii	2n=2x=22	Echinate-cristate	390–510	Echinate	22–27 μm
			(mean=450 μm)		(mean=25 μm)
I. longpingii	2n=4x=44	Tuberculate-cristate	310-410 μm	Echinate	27–30 μm
			(mean=350 μm)		(mean=29 µm)
I. xiangfei	2n=4x=44	Cristate-reticulate	390-450 μm	Tuberculate	26–28 μm
			(mean=430 μm)		(mean=27 µm)

Note: * Cited from Liu et al. 2008. Data of *I. shangrilaensis* and *I. baodongii* were cited from Li et al. (2019) and Lu et al. (2021), respectively.

3 Keys to Iso ëtes of China

(1) Individual diploid	(2)
(1) Individual polyploid	(6)
(2) Megaspore levigate or just tuberculate-regulate	(3)
(2) Megaspore with various type of ornamentation	(4)
(3) Leaves 2–5 cm long, microspores echinate	I. hypsophila
(3) Leaves 3–18 cm long, microspores echinate to cristate	I. shangrilaensis
(4) Megaspore tuberculate-scristae, just distributing in Taiwan	I. taiwanensis
(4) Megaspore echinate or cristate-reticulate	
(5) Microspore echinate, distributing in Zhejiang	
(5) Microspore levigate-granulate, distributing in Yunnan and Guizhou	
(6) Leaves ca. 1 mm wide at mid-length	(7)
(6) Leaves 2–3 mm wide at mid-length	I. xiangfei
(7) Megaspore ornamentation cristate-reticulate, chromosome number 66	I. orientalis
(7) Megaspore without reticulate ornamentation, chromosome number 44	(8)
(8) Rhizome 2-lobed, megaspore 310–410 μm (mean=350 μm)	I. longpingii
(8) Rhizome 3-lobed, megaspore 340–450 μ m (mean=409 μ m)	I. sinensis

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